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## SEQUENCE LISTING

- (1) GENERAL INFORMATION:
  - (I) APPLICANT: Bestor, Timothy H.
  - (ii) TITLE OF INVENTION: CHIMERIC DNA-BINDING/DNA METHYLTRANSFERASE NUCLEIC ACID AND POLYPEPTIDE AND USES

THEREOF

- (iii) NUMBER OF SEQUENCES: 2
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Cooper & Dunham LLP
  - (B) STREET: 1185 Avenue of the Americas
  - (C) CITY: New York
  - (D) STATE: New York
  - (E) COUNTRY: U.S.A.
  - (F) ZIP: 10036
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE: 31-JAN-1996
  - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:

  - (A) NAME: White, John P.(B) REGISTRATION NUMBER: 28,678
  - (C) REFERENCE/DOCKET NUMBER: 48075-A
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 212-278-0400
    - (B) TELEFAX: 212-391-0526
- (2) INFORMATION FOR SEQ ID NO:1:
  - (I) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 632 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA (genomic)
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:



TGGAAGGGCT	AATTCACTCC	CAACGAAGAC	AAGATATCCT	TGATCTGTGG	ATCTACCACA	60
CACAAGGCTA	CTTCCCTGAT	TAGCAGAACT	ACACACCAGG	GCCAGGAGTC	AGATATCCAC	120
TGACCTTTGG	ATGGTGCTAC	AAGCTAGTAC	CAGTTGAGCC	AGATAAGGTA	GAAGAGGCCA	180
ACAAAGGAGA	GAACACCAGC	TTGTTACACC	CTGTGAGCCT	GCATGGGATG	GATGACCCGG	240
ATATATAAGT	GTTAGAGTGG	AGGTTTGACA	GCCGCCTAGC	ATTTCATCAC	GTGGCCCGAG	300
AGCTGCATCC	GGAGTACTTC	AAGAACTGCT	GATATCGAGC	TTGCTACAAG	GGACTTTCCG	360
CTGGGGACTT	TCCAGGGAGG	CGTGGCCTGG	GCGGGACTGG	GGAGTGGCGA	GCCCTCAGAC	420
CTGCATATAA	GCAGCTGCTT	TTTGCCTGTA	CTGGGTCTCT	CTGGTTAGAC	CAGATCTGAG	480
CCTGGGAGCT	CTCTGGCTAG	CTAGGGAACC	CACTGCTTAA	GCCTCAATAA	AGCTTGCCTT	540
GAGTGCTTCA	AGTAGTGTGT	GCCCGTCTGT	TGTGTGACTC	TGGTAACTAG	AGATCCCTCA	600
GACCCTTTTA	GTCAGTGTGG	AAAATCTCTA	GC			632

## (2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 40 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
    (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: CCCCCTGTG AGCCTGCGGG TTTTTTCCCG CAGGCTCACA

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